

Figure 1.

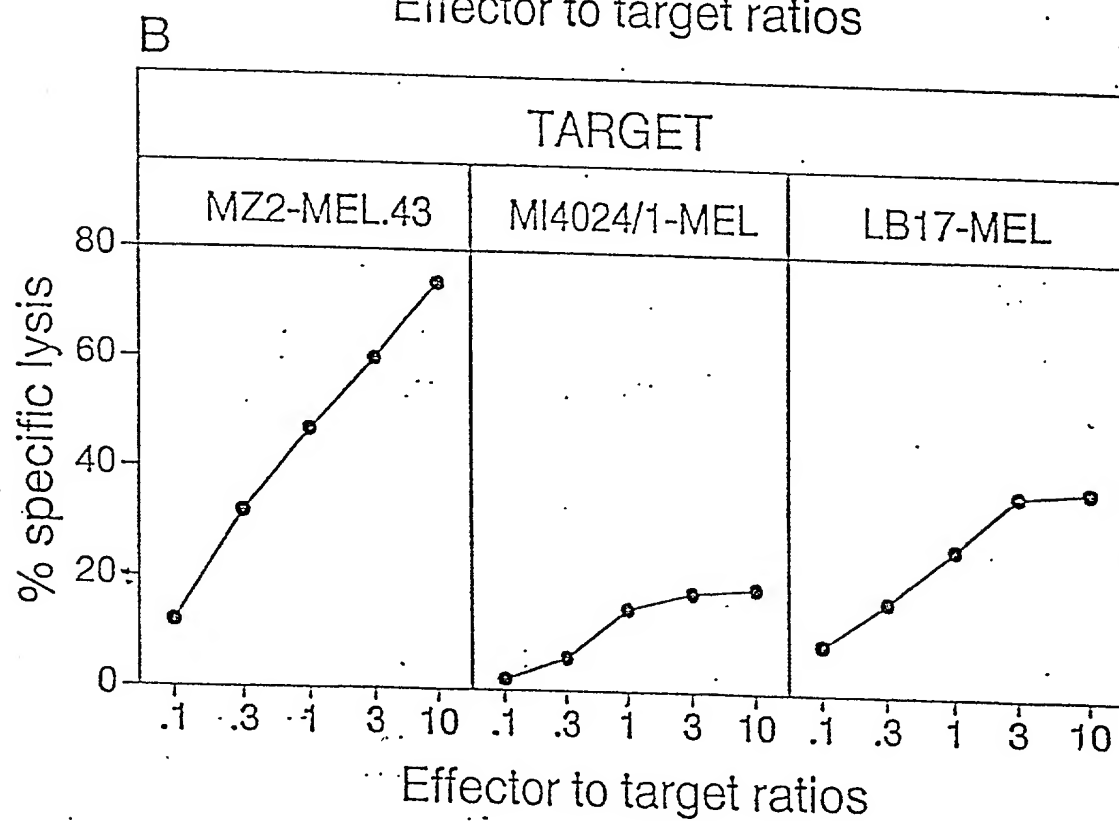
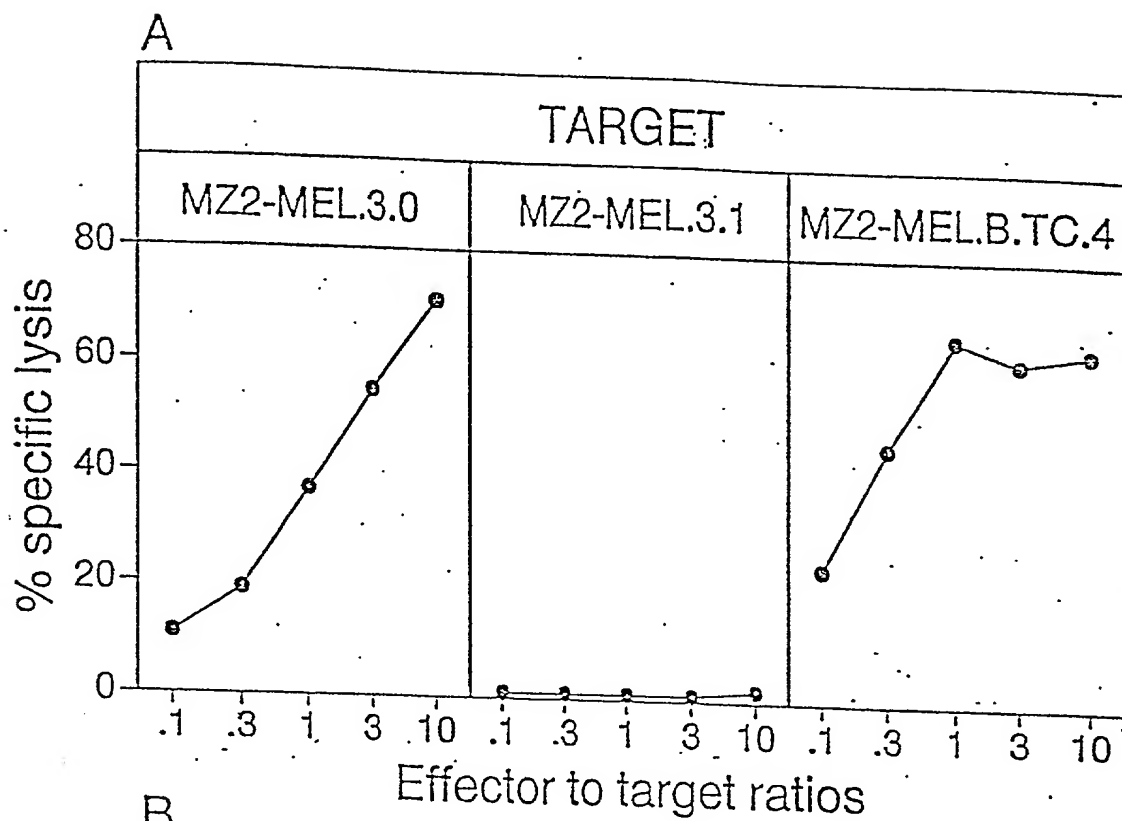


Figure 3

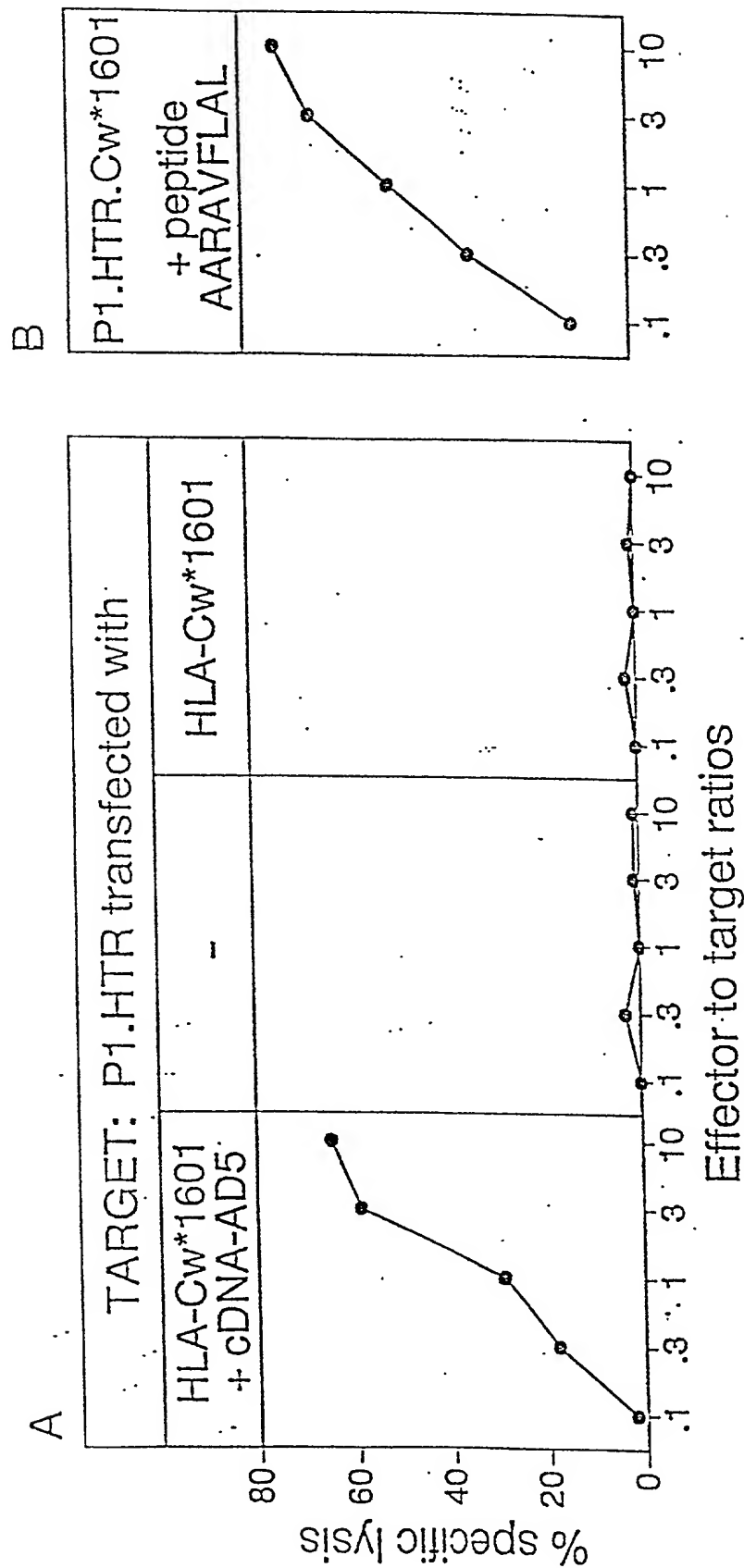


Figure 4

CGCCAATTTA	GGGTCTCCGG	TATCTCCCGC	TGAGCTGCTC	TGTTCCCGGC	TTAGAGGACC	60
AGGAGAAGGG	GGAGCTGGAG	GCTGGAGCCT	GTAACACCGT	GGCTCGTCTC	ACTCTGGATG	120
GTGGTGGCAA	CAGAGATGGC	AGCGCAGCTG	GAGTGTTAGG	AGGGCGGCCT	GAGCGGTAGG	180
		M	A A R A V F	L A L	S A Q	13
AGTGGGGCTG	GAGCAGTAAG	ATGGCCGCCA	GAGCGGTTTT	TCTGGCATTG	TCTGCCCAGC	240
L L Q A	R L M	K E E	S P V V	S W R	L E P	33
TGCTCCAAGC	CAGGCTGATG	AAGGAGGAGT	CCCCTGTGGT	GAGCTGGAGG	TTGGAGCCTG	300
E D G T	A L C	F I F				43
AAGACGGCAC	AGCTCTGTGC	TTCATCTTCT	GAGGTTGTGG	CAGCCACGGT	GATGGAGACG	360
GCAGCTCAAC	AGGAGCAATA	GGAGGAGATG	GAGTTTCACT	GTGTCAGCCA	GGATGGTCTC	420
GATCTCCTGA	CCTCGTGATC	CGCCCGCCTT	GGCCTTCCAA	AGTGCCGAGA	TTACAGCGAT	480
GTCGATTTTG	TAAGCACTTT	GGAGCCACTA	TCAAATGCTG	TGAAGAGAAA	TGTACCCAGA	540
TGTATCATT	TCCTTGTGCT	GCAGGAGCCG	GCTCCTTTCA	GGATTTCACT	CACATCTTCC	600
TGCTTTGTCC	AGAACACATT	GACCAAGCTC	CTGAAAGATG	TAAGTTTACT	ACGCATAGAC	660
TCTTAACTT	CAACCAATGT	ATTTACTGAA	AATAACAAAT	GTTGTAAATT	CCCTGAGTGT	730
TATCTTACTT	GTATTAAAAG	GTAATAATAC	ATAATCATT	AAATCTGAGG	GATCATTGCC	780
AGAGATTGTT	GGGGAGGGAA	ATGTTATCAA	CGGTTTCATT	GAAATTAAAT	CCAAAAAGTT	840
ATTCCTCAG	AAAAATCAAA	TAAAGTTTGC	ATGTTTTTTA	TTCTTAAAAC	ATTTTAAAAA	900
CCACTGTAGA	ATGATGTAAA	TAGGGACTGT	GCAGTATTTT	TGACATATAC	TATAAAATTA	960
TTAAAAAGTC	AATCAGTATT	CAACATCTTT	TACACTAAAA	AGCC		1004

Figure 5

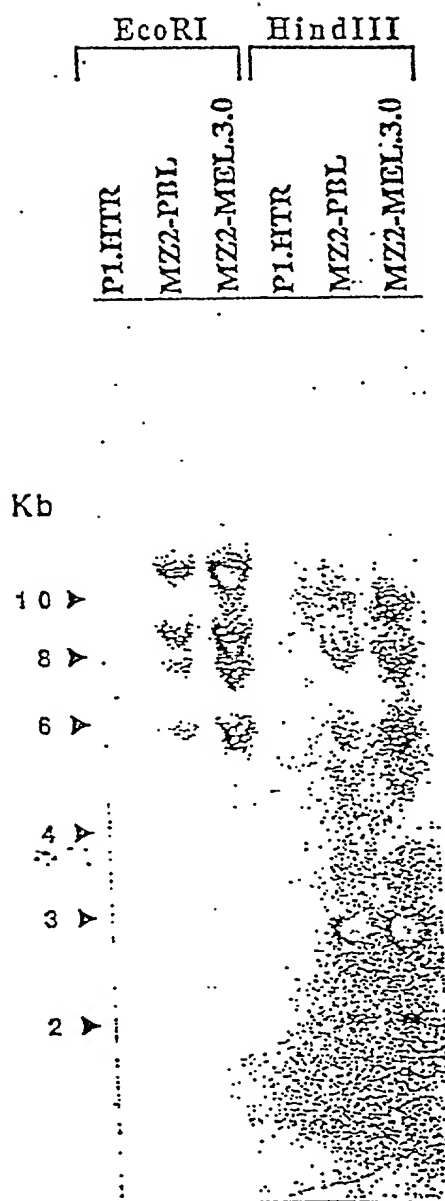


Figure 6

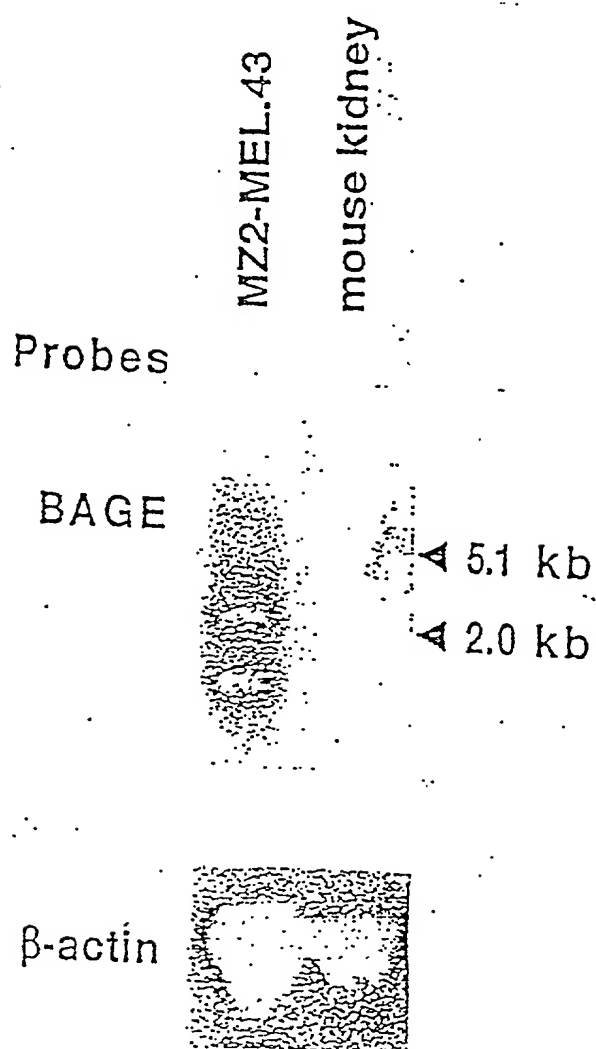


Figure 7

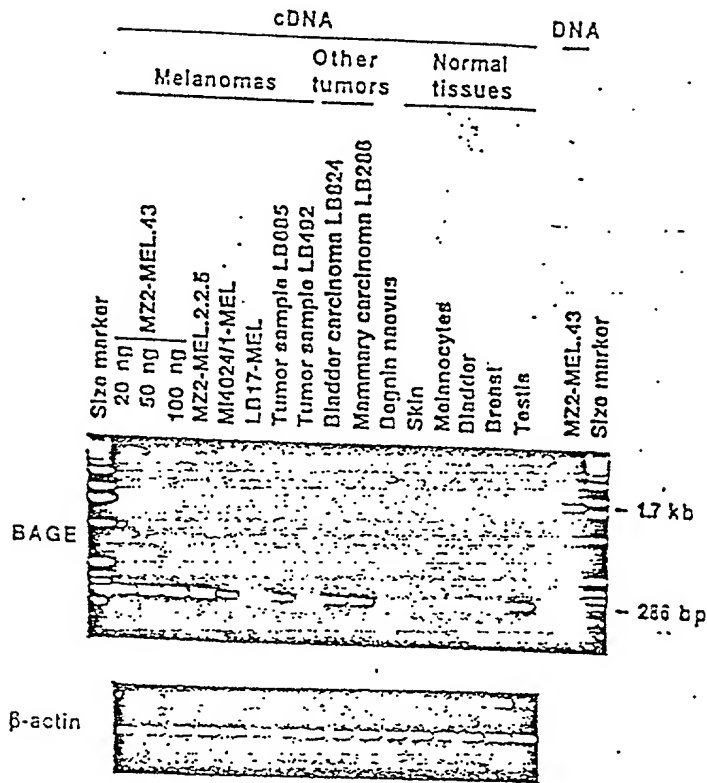


Figure 8

